

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/532, 028

Source: PCT

Date Processed by STIC: 04/11/2006

ENTERED

PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/532,028

DATE: 04/11/2006

TIME: 12:37:36

Input Set : A:\65959-50.txt

Output Set: N:\CRF4\04112006\J532028.raw

```

3 <110> APPLICANT: Boender, Piet
5 <120> TITLE OF INVENTION: IMPROVED METHODS FOR GENERATING MULTIPLE RNA COPIES
7 <130> FILE REFERENCE: 65959/50
9 <140> CURRENT APPLICATION NUMBER: 10/532,028
10 <141> CURRENT FILING DATE: 2005-04-21
12 <150> PRIOR APPLICATION NUMBER: PCT/EP03/12058
13 <151> PRIOR FILING DATE: 2003-10-30
15 <150> PRIOR APPLICATION NUMBER: US 60/440,688
16 <151> PRIOR FILING DATE: 2003-01-17
18 <150> PRIOR APPLICATION NUMBER: EP 02447204.5
19 <151> PRIOR FILING DATE: 2002-10-30
21 <160> NUMBER OF SEQ ID NOS: 3
23 <170> SOFTWARE: PatentIn version 3.3
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 55
27 <212> TYPE: DNA
28 <213> ORGANISM: Artificial Sequence
30 <220> FEATURE:
31 <223> OTHER INFORMATION: Primer
34 <220> FEATURE:
35 <221> NAME/KEY: misc_feature
36 <222> LOCATION: (50)..(55)
37 <223> OTHER INFORMATION: N = an equimolar mixture of A, C, G and T bases of which the
last
38     nucleotide is a 2'-3' dideoxynucleotide and the final phosphate
39     bond has been exchanged with a phosphorothioate bond
41 <400> SEQUENCE: 1
W--> 42 aattctaata cgactcacta tagggagaga aggataccac tagctagcgn nnnnn      55
45 <210> SEQ ID NO: 2
46 <211> LENGTH: 36
47 <212> TYPE: DNA
48 <213> ORGANISM: Artificial Sequence
50 <220> FEATURE:
51 <223> OTHER INFORMATION: Primer
54 <220> FEATURE:
55 <221> NAME/KEY: misc_feature
56 <222> LOCATION: (31)..(36)
57 <223> OTHER INFORMATION: N stands for an equimolar mixture of A, C, G and T bases - of
58     which the last nucleotide is a 2'-3' dideoxynucleotide
60 <400> SEQUENCE: 2
W--> 61 aattctaata cgactcacta tagggagaga nnnnnn      36
64 <210> SEQ ID NO: 3
65 <211> LENGTH: 30
66 <212> TYPE: DNA

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67 <213> ORGANISM: Artificial Sequence

69 <220> FEATURE:

70 <223> OTHER INFORMATION: Primer

72 <400> SEQUENCE: 3

73 agcacggaaa gacgatgacg gaaaaagaga

30

RAW SEQUENCE LISTING ERROR SUMMARY
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Input Set : A:\65959-50.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 50,51,52,53,54,55

Seq#:2; N Pos. 31,32,33,34,35,36

VERIFICATION SUMMARY

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Input Set : A:\65959-50.txt

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L:42 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0

L:61 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0